
Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2009; month=8; day=27; hr=12; min=53; sec=30; ms=213;]

Validated By CRFValidator v 1.0.3

Application No: 10578840 Version No: 2.0

Input Set:

Output Set:

Started: 2009-07-30 11:00:11.438 **Finished:** 2009-07-30 11:00:19.922

Elapsed: 0 hr(s) 0 min(s) 8 sec(s) 484 ms

Total Warnings: 122

No. of SeqIDs Defined: 122

Actual SeqID Count: 122

Total Errors:

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W	213	Artificial or Unknown found in <213> in SEQ ID (2))
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W	213	Artificial or Unknown found in <213> in SEQ ID (4))
W	213	Artificial or Unknown found in <213> in SEQ ID (5))
W	213	Artificial or Unknown found in <213> in SEQ ID (6))
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W	213	Artificial or Unknown found in <213> in SEQ ID (9))
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Input Set:

Output Set:

Started: 2009-07-30 11:00:11.438 **Finished:** 2009-07-30 11:00:19.922

Elapsed: 0 hr(s) 0 min(s) 8 sec(s) 484 ms

Total Warnings: 122
Total Errors: 2

No. of SeqIDs Defined: 122
Actual SeqID Count: 122

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SEQUENCE LISTING

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133

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                                    -10
                                                                       96
gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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                                                                      144
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Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt
                                                                      192
Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
30
                    35
                                         40
                                                             45
gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat
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Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
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                                    55
gag aag ttc aag gac aga gtc acg atg acc cgg gac acg tcc acg agc
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Glu Lys Phe Lys Asp Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser
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                                                     75
aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc
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Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
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tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa
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Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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424

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                -15
                                                         -5
                                     -10
gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
                                                                        96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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Gly Thr Leu Val Thr Val Ser Ser

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gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu 30 35 40 45	192										
gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn 50 55 60	240										
gag aag ttc aag gac aga gtc acg atg acc tca gac acg tcc acg agc Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Thr Ser 65 70 75	288										
aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val 80 85 90	336										
tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln 95 100 105	384										
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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
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Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc
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Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
    15
                        20
acc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt
                                                                      192
Thr Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
                    35
                                                                      240
gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
                50
gag aag ttc aag gac aga gtc acg atg acc tca gac acg tcc acg agc
                                                                      288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Thr Ser
            65
                                70
                                                    75
aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc
                                                                      336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
        80
                            85
                                                                      384
tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
    95
                        100
                                            105
                                                                      424
gga acc ctg gtc acc gtc tcc tca ggtgagtgga tccgcg
Gly Thr Leu Val Thr Val Ser Ser
110
                  115
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                -15
                                    -10
                                                         -5
gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
                                                                       96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc
                                                                      144
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Pro	Gly 15	Ala	Ser	Val	Lys	Val 20	Ser	Cys	Lys	Ala	Ser 25	Gly	Tyr	Thr	Phe	
-			-					cga Arg	_	-						192
								tac Tyr								240
	_		_	_		-	_	atg Met 70			_	_		_	_	288
	-		_		-	-	-	ctc Leu	_			-	_	-	-	336
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		-	gtc Val		-			ggto	gagto	gga t	ccgo	cg				424
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)> 17 aagga		gagto	cacgo	ct ga	accto	agad	c acç	gtcca	acg						39
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                -15
                                    -10
                                                                        96
gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
        -1 1
                                                 10
cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc
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Glu	Trp	Met	Gly	Tyr	Ile	Tyr	Pro	Tyr	Asn	Asp	Gly	Thr	Lys	Tyr	Asn	
	_		_	50		_		_	55	_	_		_	60		
gag	aad	ttc	aad	gac	aga	atc	aca	atq	acc	tca	gac	acq	tac	acq	age	288
gag	aag	ttc	aag	gac	aga	gtc	acg	atg	acc	tca	gac	acg	tcc	acg	agc	288
	aag Lys		_	-	-	-	_	_			_	_		_	-	288
	_		_	-	-	-	_	_			_	_		_	-	288

aca gtc